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		Application Number	
		Filing Date	
		First Named Inventor	Adam T. Zemla
		Group Art Unit	
		Examiner Name	
Sheet 1 of 2	Attorney Docket Number	IL-11160	

U.S. PATENT DOCUMENTS						
Examiner Initials*	Cite No. <sup>1</sup>	U.S. Patent Document		Name of Patentee or Applicant of Cited Document	Date of Publication of Cited Document MM-DD-YYYY	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear
		Number	Kind Code <sup>2</sup> (if known)			
MW		2002/0150906	A1	Debe	10-17-2002	
MW		2003/0130797	A1	Skolnick, et al.	07-10-2003	

FOREIGN PATENT DOCUMENTS							
Examiner Initials*	Cite No. <sup>1</sup>	Office <sup>3</sup>	Foreign Patent Document		Name of Patentee or Applicant of Cited Document	Date of Publication of Cited Document MM-DD-YYYY	Pages, Columns, Lines, Where Relevant Passages or Relevant Figures Appear
			Number <sup>4</sup>	Kind Code <sup>5</sup> (if known)			
MW			WO 93/01484		Eisenberg, et al.	01-21-1993	
			WO 98/48270		Goddard, et al.	10-29-1998	
MW			WO 00/11206		Skolnick, et al.	03-02-2000	

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Sheet 2 of 2	Attorney Docket Number		IL-11160
<b>OTHER PRIOR ART - NON PATENT LITERATURE DOCUMENTS</b>			
Examiner Initials*	Cite No. <sup>1</sup>	Include name of the author (in CAPITAL LETTERS), title of the article (when appropriate), title of the item (book, magazine, journal, serial, symposium, catalog, etc.), date, page(s), volume-issue number(s), publisher, city and/or country where published.	T 2
MW		ZEMLA, A., "LGA: a method for finding 3D similarities in protein structures," Nucleic Acids Research, Oxford University Press, Vol. 31, No. 13, 2003, pp. 3370-3374	
		ZU-KANG, F., et al., "Optimum superimposition of protein structures: ambiguities and implications," Research Paper, Folding & Design, Center for Applied Molecular Engineering, Institute for Chemistry and Biochemistry, University of Salzburg, Austria, Feb. 26, 1996, pp.123-132	
		LACKNER, P., et al., "ProSup: a refined tool for protein structure alignment," Protein Engineering, Oxford University Press, Vol. 13, No. 11, 2000, pp. 745-752	
		SHINDYALOV, I., et al., "Protein structure alignment by incremental combinatorial extension (CE) of the optimal path," Protein Engineering, Vol. 11, No. 9, 1998, pp. 739-747	
MW		HOLM, L., et al., "Protein Structure Comparison by Alignment of Distance Matrices," J. Mol. Biol., Vol. 233, 1993, pp. 123-138	

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